FIG. 1

```
ATGCCCAAGCGCGCGCACTGGGGGGCCCTCTCCGTGGTGCTGATCCTGCTTTGGGGCCATCCGCGAGTGGCGCTGGCCTGCCCGCATCCTTGTGCCT
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GCTTTAAGAGACCTCAGCTCTCTTCAGGTTTTCAAGTTCAGCTACAACAAGCTGAGAGTGATCACAGGACAGACCCTCCAGGGTCTCTCTAACTTAA
TGAGGCTGCACATTGACCACAACAAGATCGAGTTTATCCACCCTCAAGCTTTCAACGTCTCTGAGGCTACTCCATTTGGAAGGAAATCT
CCTCCACCAGCTGCACCCCAGCACCTTCTCCACGTTCACATTTTTGGATTATTTCAGACTCTCCACCATAAGGCACCTCTACTTAGCAGAGAACATG
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GGTTTTTGGAATGGGATGCAAAATCCAGAGGAATTCTGAAGTGTAAAAAGGACAAAGCTTATGAAGGCGGTCAGTTGTGTGCAATGTGCTTCAGTCC
AAAGAAGTTGTACAAACATGAGATACACAAGCTGAAGGACATGACTTGTCTGAAGCCTTCAATAGAGTCCCCTCTGAGACAGAACAGGAGCAGGAGT
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CATAAATGCAACAGTTGCCTTGGACTTTGAGTGTCCAATGACCCGAGAAAACTATGAAAAGCTATGGAAATTGATAGCATACTACAGTGAAGTTCCC
GTGAAGCTACACAGAGAGCTCATGCTCAGCAAAGACCCCAGAGTCAGCTACCAGTACAGGCAGGATGCTGATGAGGAAGCTCTTTACTACACAGGTG
TGAGAGCCCAGATTCTTGCAGAACCAGAATGGGTCATGCAGCCATCCATAGATATCCAGCTGAACCGACGTCAGAGTACGGCCAAGAAGGTGCTACT
TTCCTACTACACCCAGTATTCTCAAACAATATCCACCAAAGATACAAGGCAGGCTCGGGGCAGAAGCTGGGTAATGATTGAGCCTAGTGGAGCTGTG
CAAAGAGATCAGACTGTCCTGGAAGGGGGTCCATGCCAGTTGAGCTGCAACGTGAAAGCTTCTGAGAGTCCATCTATCTTCTGGGTGCTTCCAGATG
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CTGTACATTTGATAATAAAAATAATATTCTCCCAAAAAAAAA

FIG. 2

MPKRAHWGALSVVLILLWGHPRVALACPHPCACYVPSEVHCTFRSLASVPAGIARHVERINLGFNSIQALSETSFAGLTKLELLMIHGNEIPSIPDG
ALRDLSSLQVFKFSYNKLRVITGQTLQGLSNLMRLHIDHNKIEFIHPQAFNGLTSLRLLHLEGNLLHQLHPSTFSTFTFLDYFRLSTIRHLYLAENM
VRTLPASMLRNMPLLENLYLQGNPWTCDCEMRWFLEWDAKSRGILKCKKDKAYEGGQLCAMCFSPKKLYKHEIHKLKDMTCLKPSIESPLRQNRSRS
IEEEQEQEEDGGSQLILEKFQLPQWSISLNMTDEHGNMVNLVCDIKKPMDVYKIHLNQTDPPDIDINATVALDFECPMTRENYEKLWKLIAYYSEVP
VKLHRELMLSKDPRVSYQYRQDADEEALYYTGVRAQILAEPEWVMQPSIDIQLNRRQSTAKKVLLSYYTQYSQTISTKDTRQARGRSWVMIEPSGAV
QRDQTVLEGGPCQLSCNVKASESPSIFWVLPDGSILKAPMDDPDSKFSILSSGWLRIKSMEPSDSGLYQCIAVRDEMDRMVYRVLVQSPSTQPAEK
DTVTIGKNPGESVTLPCNALAIPEAHLSWILPNRRIINDLANTSHVYMLPNGTLSIPKVQVSDSGYYRCVAVNQQGADHFTVGITVTKKGSGLPSKR
GRRPGAKALSRVREDIVEDEGGSGMGDEENTSRRLLHPKDQEVFLKTKDDAINGDKKAKKGRRKLKLWKHSEKEPETNVAEGRRVFESRRRINMANK
QINPERWADILAKVRGKNLPKGTEVPPLIKTTSPPSLSLEVTPPFPAVSPPSASPVQTVTSAEESSADVPLLGEEEHVLGTISSASMGLEHNHNGVI
LVEPEVTSTPLEEVVDDLSEKTEEITSTEGDLKGTAAPTLISEPYEPSPTLHTLDTVYEKPTHEETATEGWSAADVGSSPEPTSSEYEPPLDAVSLA
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ENKHRNIVTPSSETILLPRTVSLKTEGPYDSLDYMTTTRKIYSSYPKVQETLPVTYKPTSDGKEIKDDVATNVDKHKSDILVTGESITNAIPTSRSL
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ETTTLDQDHLETTVAILLSETRPQNHTPTAARMKEPASSSPSTILMSLGQTTTTKPALPSPRISQASRDSKENVFLNYVGNPETEATPVNNEGTQHM
SGPNELSTPSSDRDAFNLSTKLELEKQVFGSRSLPRGPDSQRQDGRVHASHQLTRVPAKPILPTATVRLPEMSTQSASRYFVTSQSPRHWTNKPEIT
TYPSGALPENKQFTTPRLSSTTIPLPLHMSKPSIPSKFTDRRTDQFNGYSKVFGNNNIPEARNPVGKPPSPRIPHYSNGRLPFFTNKTLSFPQLGVT
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VLLSVTVQQPQILASHYQDVTVYLGDTIAMECLAKGTPAPQISWIFPDRRVWQTVSPVESRITLHENRTLSIKEASFSDRGVYKCVASNAAGADSLA
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VQLNVQRAAANARITGTSPRRTDVRYGGTLKLDCSASGDPWPRILWRLPSKRMIDALFSFDSRIKVFANGTLVVKSVTDKDAGDYLCVARNKVGDDY
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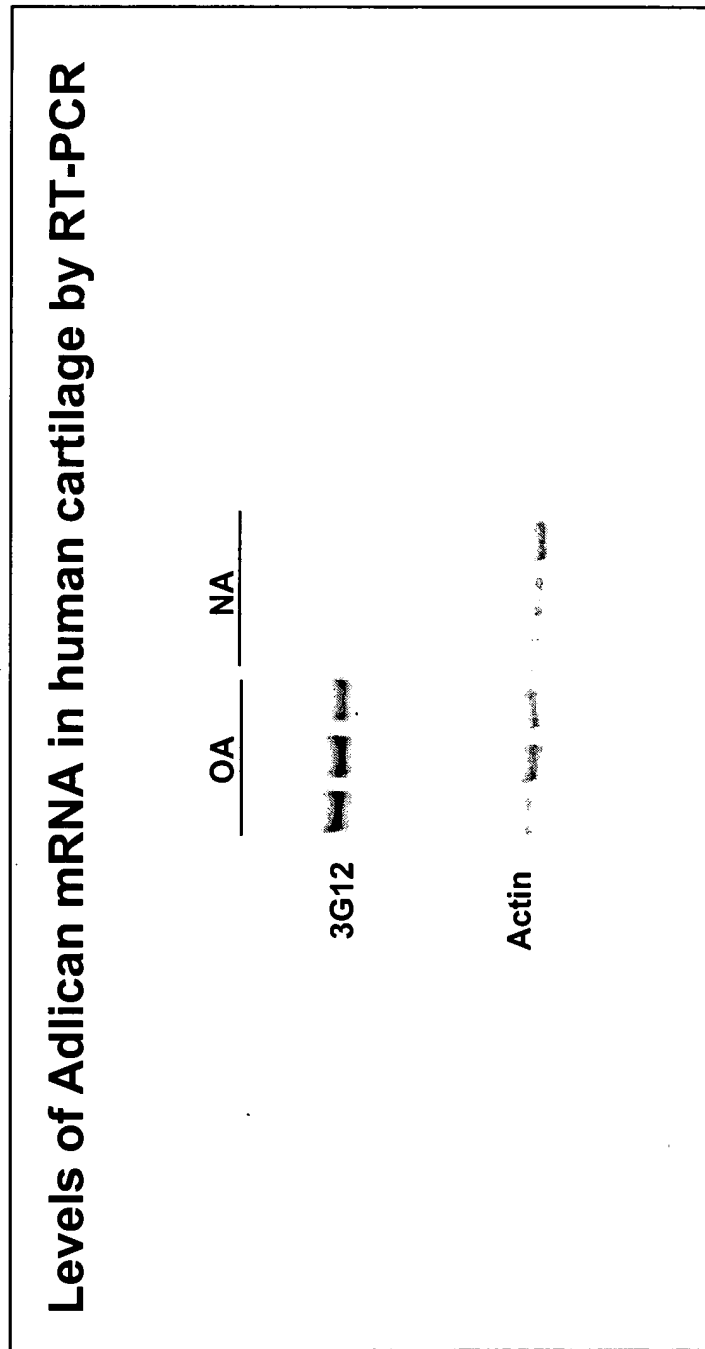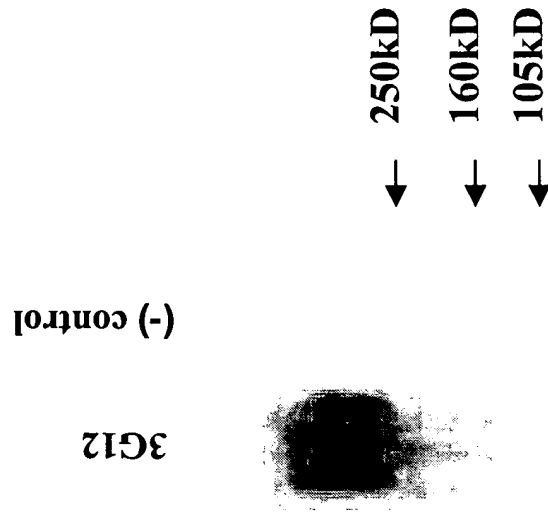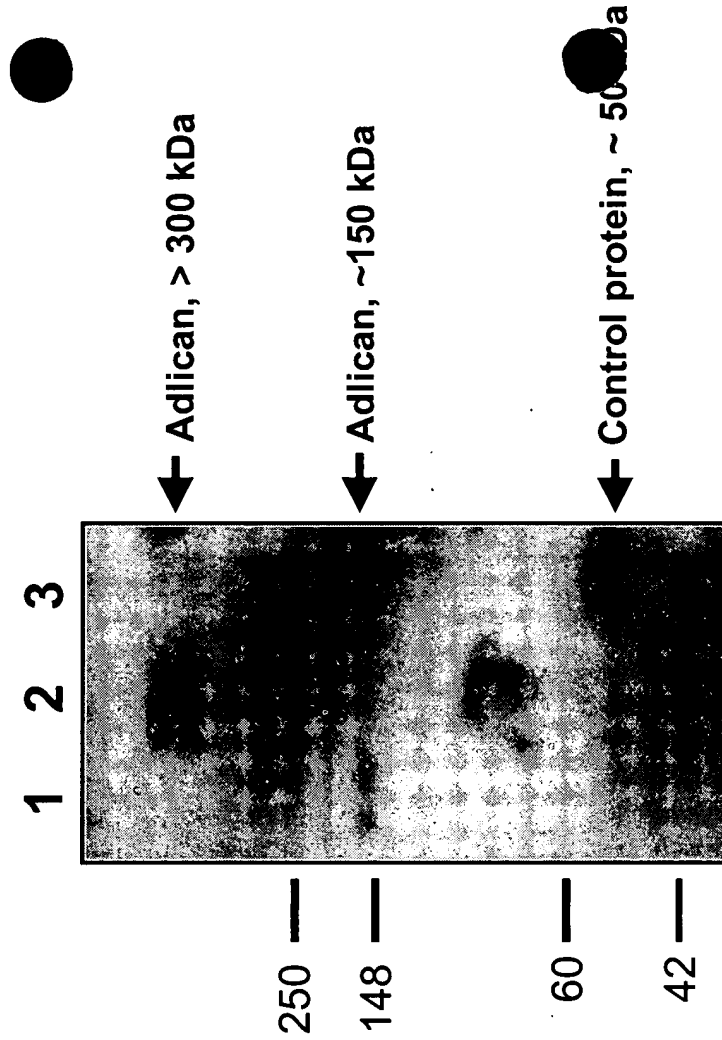NILGSDSKTTYIHVF

FIG. 3

Levels of Adlican mRNA in human cartilage by RT-PCR

OA   NA

3G12

Actin

FIG. 5

Adlican, > 300 kDa

Adlican, ~150 kDa

Control protein, ~ 50 kDa

1 2 3

250 —

148 —

60 —

42 —

FIG. 4

3G12

(-) control

250kD

160kD

105kD

FIG. 6

Patient: trauma | OA | RA

04 40   11 18 34 13   30 41 42

kDa
205—
139—
84—
45—

DIAGNOSIS	Western blot positive/total
trauma	1/2
gout	0/3
OA, mild/mod.	2/4
OA, severe	4/4 ←
RA, moderate	2/6
RA, severe	1/2